

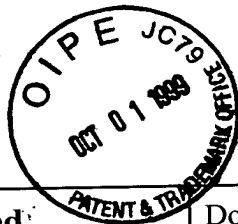


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Form PTO-1449 Modified List of Patent and Publications Cited by Applicant (Use several sheets if necessary) U.S. Department of Commerce Patent and Trademark Office		Docket No. IBIS-0012	Serial No. 09/310,667
		Applicant David J. Ecker et al.	
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	AB	Benson et al., "GenBank", <i>Nucl. Acids Res.</i> , 1998 , 26(1), 1-7	
	AC	Brown, J.W., "Phylogenetic analysis of RNA structure on the Macintosh computer", <i>CABIOS Commun.</i> , 1991 , 7(3), 391-393	
	AD	Gautheret et al., "Inferring the conformation of RNA base pairs and triples from patterns of sequence variation", <i>Nucl. Acids Res.</i> , 1997 , 25(8), 1559-1564	
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	AI	Gutell, "Collection of small subunit (16S- and 16S-like) ribosomal RNA structures", <i>Nucl. Acids Res.</i> , 1993 , 21(13), 3051-3054	
	AJ	Gutell, "A compilation of large subunit (23S and 23S-like) ribosomal RNA structures: 1993", <i>Nucl. Acids Res.</i> , 1993 , 21(13), 3055-3074	
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	AL	Laferriere et al., "An RNA pattern matching program with enhanced performance and portability", <i>Comput. Appl. Biosci.</i> , 1994 , 10(2), 211-212	
<i>u</i>	AM	Lodmell et al., "Genetic and comparative analyses reveal an alternative secondary structure in the region of nt 912 <i>Escherichia coli</i> 16S rRNA", <i>Proc. Natl. Acad. Sci. USA</i> , 1995 , 92, 10555-10559	
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*A copy of this reference will not be forwarded to the U.S. Patent and Trademark Office since it is believed to be too voluminous and easily obtainable by the Examiner.

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Patent and Trademark Office

Docket No.
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Serial No.
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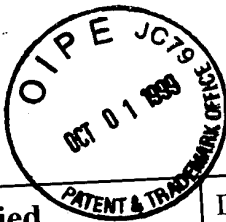
Applicant
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<i>22</i>	AN	Sutton et al., "TIGR Assembler: A New Tool for Assembling Large Shotgun Sequencing Projects", <i>Genome Science & Tech.</i> , 1995 , 1(1), 9-19
<input checked="" type="checkbox"/>	AO	Thompson et al., "CLUSTAL W: improving the sensitivity of progressive multiple sequence alignment through sequence weighting, position-specific gap penalties and weight matrix choice", <i>Nucl. Acids Res.</i> , 1994 , 22(22), 4673-4680
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	AR	Casey et al., "Iron-Responsive Elements: Regulatory RNA Sequences that Control mRNA Levels and Translation," <i>Science</i> , 1988 , 240, 924-928
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	AU	Griffey et al., "Detection of base pair mismatches in duplex DNA and RNA oligonucleotides using electrospray mass spectrometry", <i>Proc. SPIE-Int. Soc. Opt. Eng.</i> , 1997 , 2985, 82-86
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	AW	Owen et al., "Noncoding 3' sequences of the transferrin receptor gene are required for mRNA regulation by iron," <i>EMBO J.</i> , 1987 , 6(5), 1287-1293
	AX	Son, "The Structure and Regulation of Histone Genes," <i>Saenghwahak Nyusu</i> , 1993 , 13(2), 64-70
<input checked="" type="checkbox"/>	AY	Zehner et al., "RNA-protein interactions within the 3' untranslated region of vimentin mRNA," <i>Nucl. Acids Res.</i> , 1997 , 25(16), 3362-3370
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May 12, 1999

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1643**1653****U. S. PATENT DOCUMENTS**

Examiner Initial		Document No.	Date	Name	Class	Subclass
* <i>W</i>	AZ	09/076,206	05/12/98	Crooke et al.		
*	BA	09/076,214	05/12/98	Griffey et al.		
*	BB	09/076,404	05/12/98	Ecker et al.		
*	BC	09/076,405	05/12/98	Griffey et al.		
*	BD	09/076,440	05/12/98	Ecker et al.		
*	BE	09/076,447	05/12/98	Griffey et al.		

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					YES	NO
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*Pursuant to 37 C.F.R. 1.98(a)(2)(iii), no copy of a U.S. patent application need be included with an Information Disclosure Statement filed under 37 C.F.R. 1.97.

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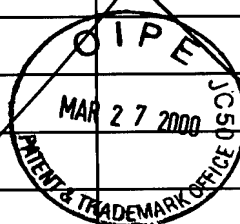
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